

FIG. 1

CG ACGGCCGGC TGGTAAATTC CCCTTCCTCC

-690 AAAATGTAAA ATAAATCTGC TTCCATCTTC TAAAATACTA TGGGACTAAA
-640 CATCCTTTG TTATGCTAAG GAAAAGCCAG TATTCGCGTT GATTTAGAAG
-590 AGGGATGTTC TGGTTATAGA ACGATGCTGT GTCTCAGAAA CACTTAAATA
-540 CTATTAAGCT AGAAATAGAA GGGAAAATAA TGCTTCCCCG CATCTCCCCT
-490 CAAGTGTAGT CCTCTTTTT TAGCCTGATT TCCGACGAAA TGTCTGAATG
-440 CCTACAGTTA TTTGGCCATC CTGAAAAGTG CAACTTATCC TGACGTCTCG

CRE

-390 AGGGACGGAA AAGTTACCGA AGTCCAAGGA ATGAGTCACT TTGCTCAAAT
-340 TTGATGAGTA ATATCAGGTG TCATGAAACC CAGTTTCGAA GGAGAGGGGA
-290 GGGGGCGTCA GATCTGCAGA CGGAAGCAGG CCGCTCCGGA TTGGATGGCG
-240 AGACCTCGAT TTTCTAAAAA TTGCGTCATT TAGAACCCAA TTGGGTCCAG

CRE-like

-190 ATGTTATGGG CATCGACGAG TTACCGTCTC GGAAACTCTC AATCACGCAA
-140 GCGAAAGGAG AGGAGGGCGC TAATTAAATA TTGAGCAGAA AGTCGCGTGG
-90 GGAGAATGTC ACGTGGGTCT GGAGGCTCAA GGAGGCTGGG ATAAATACCG
-40 CAAGGCACTG AGCAGGGCGAA AGAGCGCGCT CGGACCTCCT
+1 TTCCCGGCGG CAGCTACCGA GAGTGCAGGAG CGACCAGCGT GCGCTCGGAG

Exon 1

+51 AACCAGAGAA CTCAGCACCC CGCGGGACTG TCCGTCGCAG TAAGTGCCCG

Intron 1

+101 CGCGGTGCTG GCCCGGGCTG CCCGGGTCAAT CCCACCCCGC ATCTGTCCGA
+151 GGTGGCCGCG CTGGGGCGC CGCTGCAGCG AGGGACAGTG GGGAGACTGG
+201 CTTCCCAAAC CCCAACCCCC CTCTTGTCT TCCACCTGCA GAGTTTCCTG
+251 GTTTGAAGGT GTGGGTTGGT GGGTTAGGGG GCTGGGGGAG CTGGGATTC
+301 GGGAGAAGAG GGTTGGAGAA TCTTTGGGAC GCGATTCTCT CGCCTAACCG
+351 GTACAGGTGA GACTTCAGTC CTTATGTTTG TGATCTTGGT TCATCCGTTG
+401 TGGGGCAGAA AATTCTGTTG CTTAACTCT TGGATAACCA CCCCTAACAG
+451 ATACATTATT TCTCTTTG GTGTCTTCTC CTCCTACCC TTCCCAGAAA

Exon 2

+501 TCCGAC

FIG. 2

EM No. EV254991105US

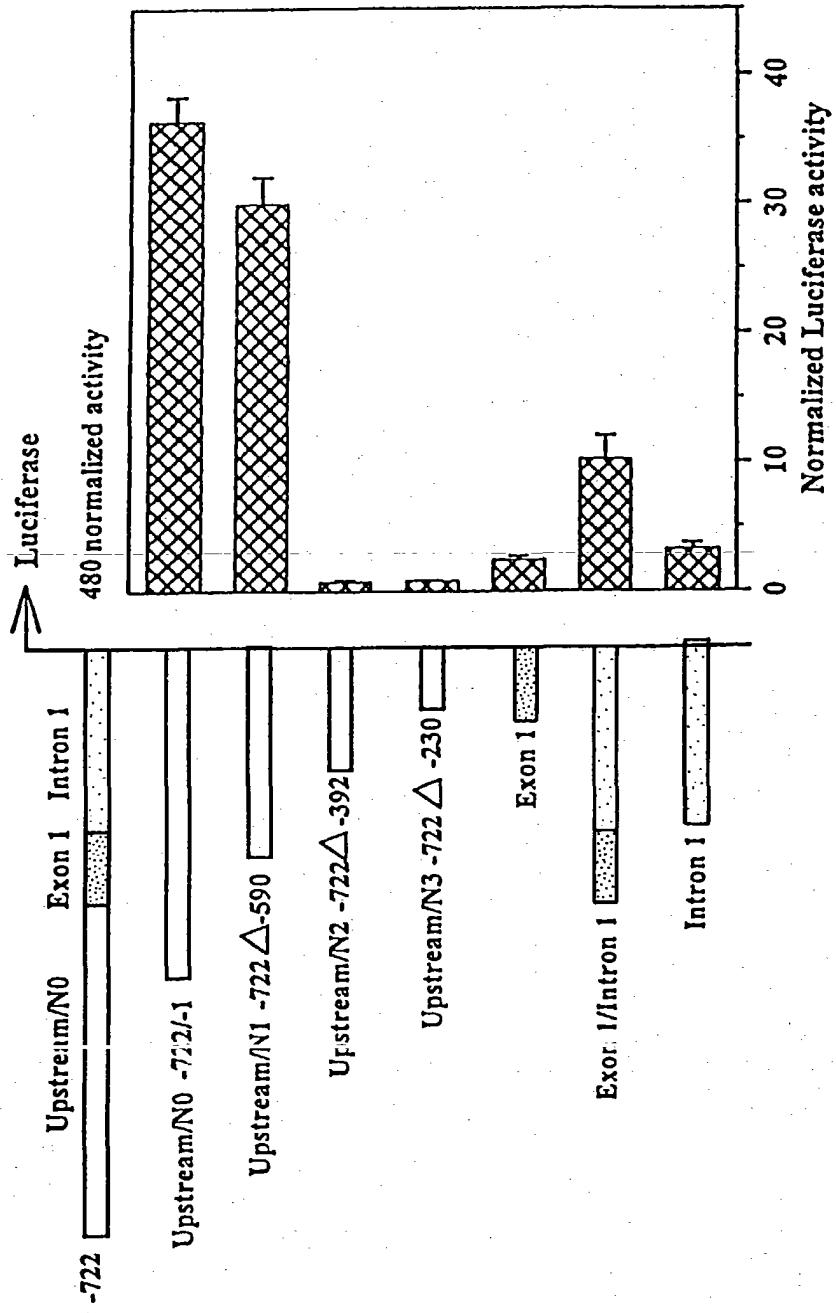
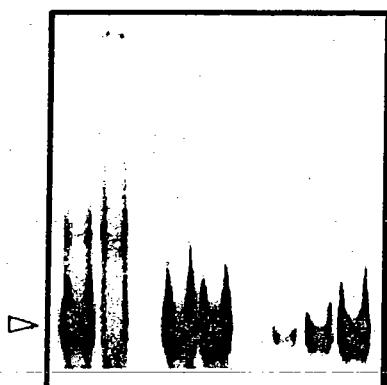
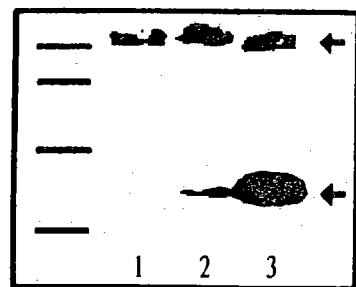


FIG. 3

FIG. 4A

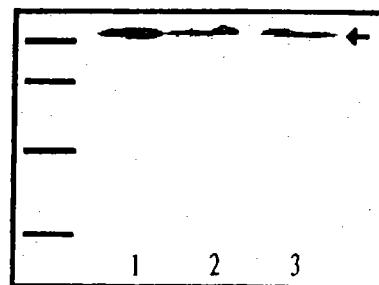


1 2 3 4 5 6 7



1 2 3

FIG. 4C



1 2 3

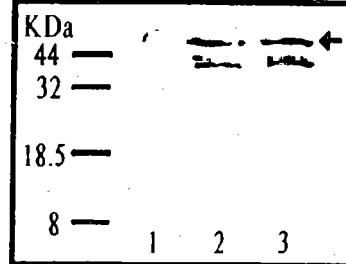


FIG. 4D

FIG. 4B

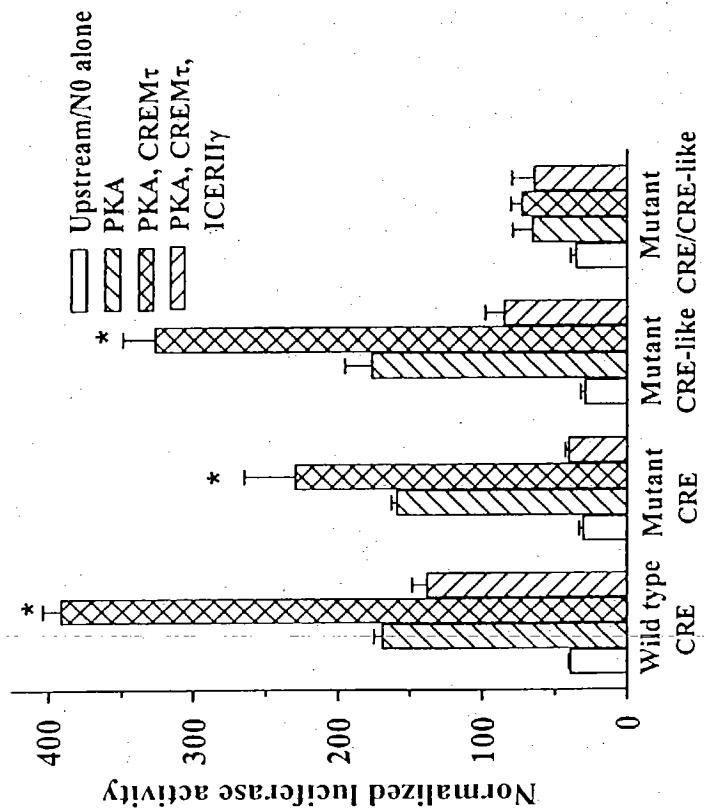


FIG. 5A

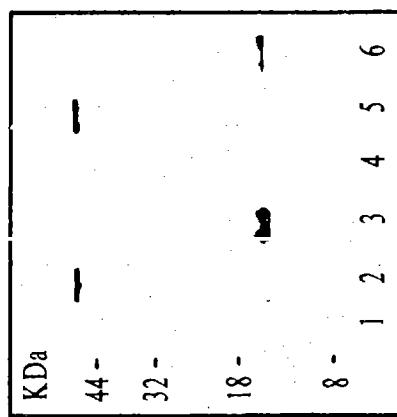


FIG. 5B

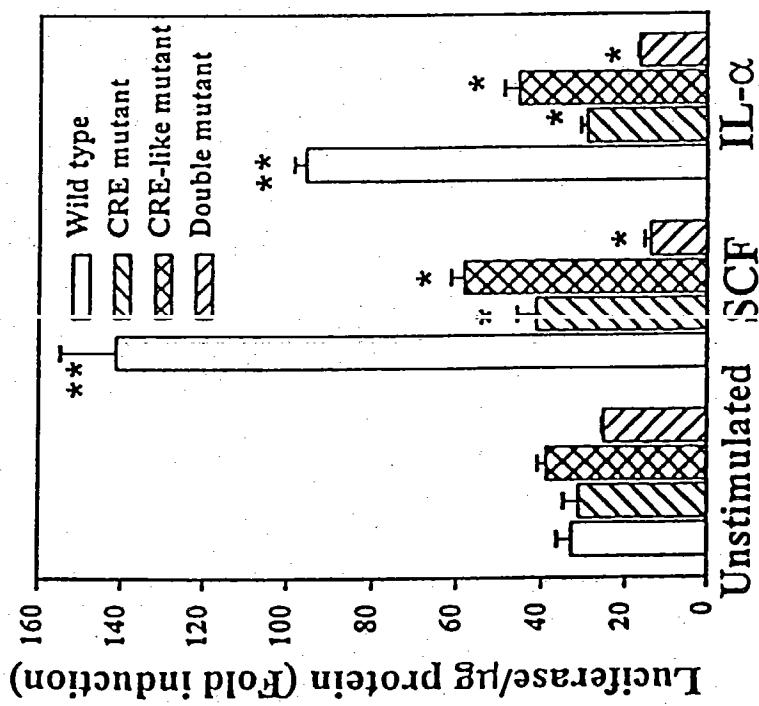


FIG. 6A

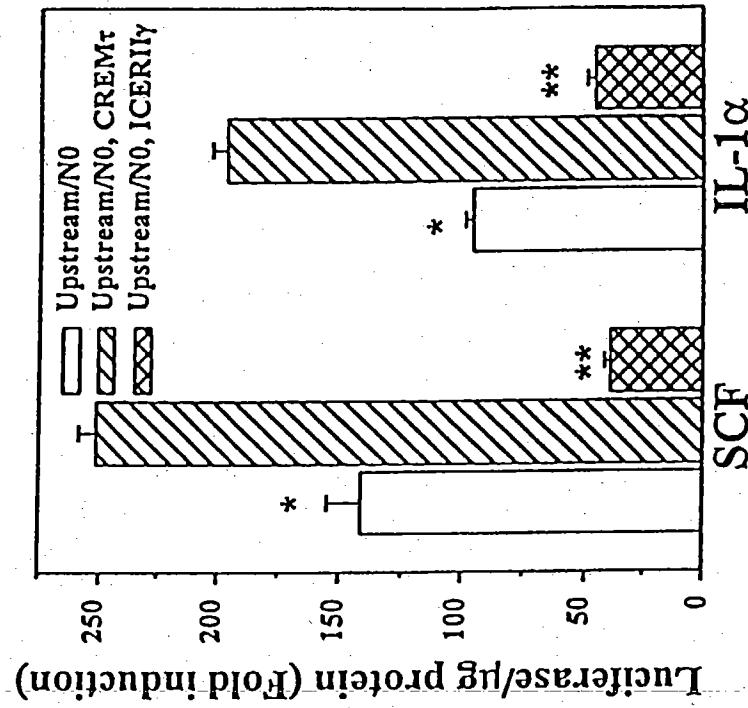


FIG. 6B

EM No. EV254991105US

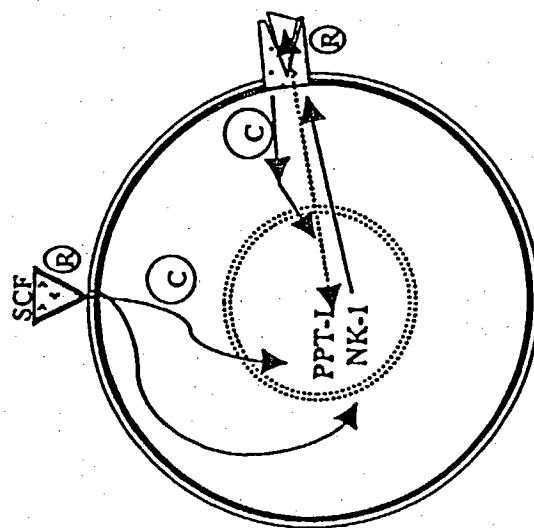


FIG. 7B

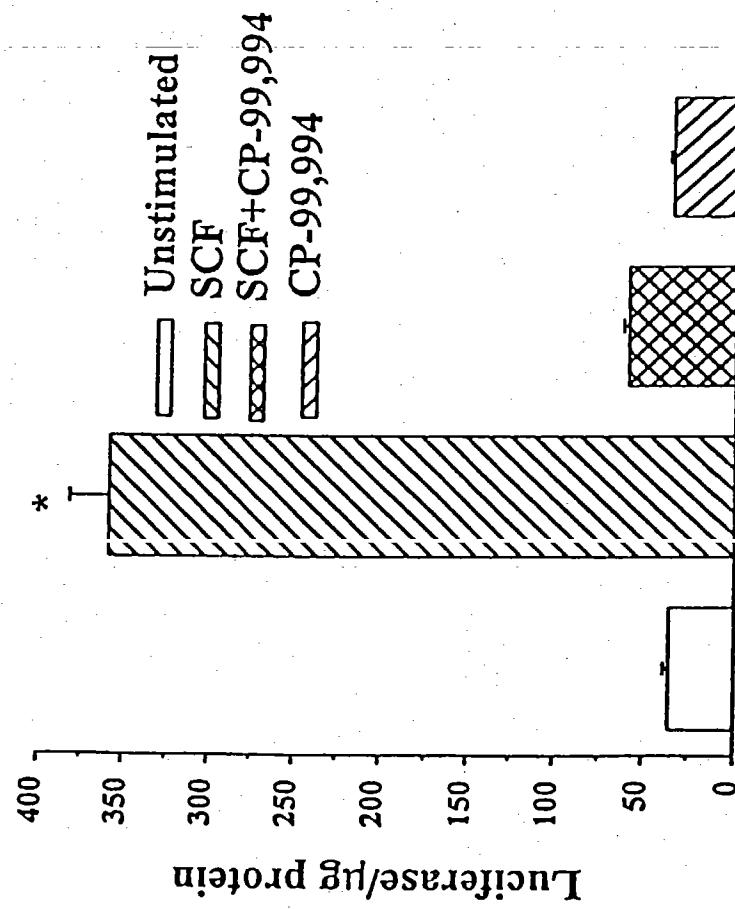


FIG. 7A

EM No. EV254991105US

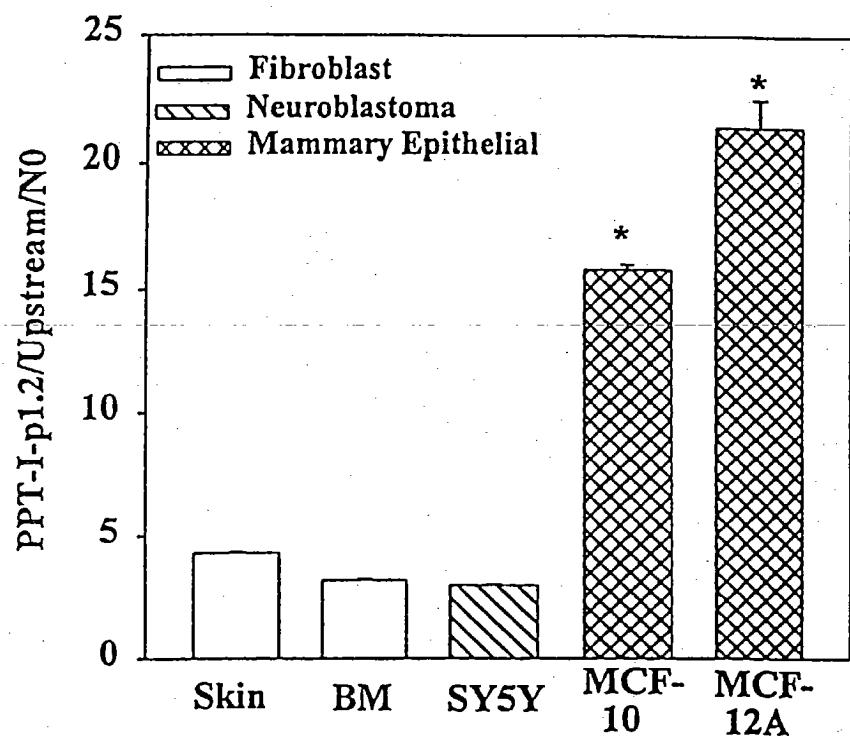


FIG. 8